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(FILE 'HOME' ENTERED AT 15:50:06 ON 16 JAN 2009)

FILE 'MEDLINE, EMBASE, BIOSIS, SCISEARCH, DISSABS, TOXCENTER' ENTERED AT
15:51:09 ON 16 JAN 2009

L1 2729 S (HEINRICH M?/AU) OR (CORLESS C?/AU)
L2 10611 S (FLETCHER J?/AU) OR (DEMETRI G?/AU)
L3 13149 S L1 OR L2
L4 218 S L3 AND (PDGF?)
L5 0 S L3 AND CD140?
L6 180 S L3 AND (PLATELET(1A) DERIVED)
L7 264 S L4 OR L6
L8 207 S L7 AND MUTAT?
L9 178 S L8 AND (TUMOR? OR CANCER? OR CARCIN?)
L10 109 S L8 AND NEOPLAS?
L11 186 S L9 OR L10
L12 456 S L3 AND GIST
L13 128 S L4 AND GIST
L14 78 DUP REM L11 (108 DUPLICATES REMOVED)
L15 7 S L14 AND PY<=2002
L16 199 DUP REM L12 (257 DUPLICATES REMOVED)
L17 26 S L16 AND PY<=2002
L18 31 S L15 OR L17
L19 31 DUP REM L18 (0 DUPLICATES REMOVED)
L20 2371 S (PDGFRA)
L21 99 S (PDGFR-A)
L22 2344 S (PDGFR(2A)ALPHA)
L23 327 S (PDGF-R(2A)ALPHA)
L24 2748 S (PDGF(2A)RECEPTOR(2A)ALPHA)
L25 4759 S (PLATELET(2A) DERIVED(5A)RECEPTOR(10A)ALPHA)
L26 23 S CD140A
L27 8628 S L20 OR L21 OR L22 OR L23 OR L24 OR L25 OR L26
L28 2386 S L27 AND MUTAT?
L29 0 S L27 AND (2072 OR 2073 OR 2074 OR 2075)
L30 1 S L27 AND (2076 OR 2077 OR 2078 OR 2079 OR 2080)
L31 0 S L27 AND (2081 OR 2082 OR 2083 OR 2084 OR 2085)
L32 0 S L27 AND (2086 OR 2087 OR 2088 OR 2089 OR 2090)
L33 0 S L27 AND (2091 OR 2092 OR 2093 OR 2094 OR 2095)
L34 0 S L27 AND (2096 OR 2097 OR 2098 OR 2099 OR 2100)
L35 0 S L27 AND (2101 OR 2102 OR 2103 OR 2104 OR 2105 OR 2106 OR 210
L36 0 S L27 AND (2916 OR 2917 OR 2918 OR 2919 OR 2920 OR 2921 OR 292
L37 0 S L27 AND (2923 OR 2924 OR 2925 OR 2926 OR 2927 OR 2928 OR 292
L38 0 S L27 AND (2930 OR 2931 OR 2932 OR 2933 OR 2934 OR 2935 OR 293
L39 39 S L27 AND (560 OR 561 OR 562 OR 563 OR 564 OR 565)
L40 156 S L27 AND (566 OR 567 OR 568 OR 569 OR 570 OR 571)
L41 186 S L39 OR L40
L42 117 DUP REM L41 (69 DUPLICATES REMOVED)
L43 48 S L27 AND (841 OR 842 OR 843 OR 844 OR 845 OR 846 OR 847 OR 84
L44 161 S L42 OR L43
L45 127 DUP REM L44 (34 DUPLICATES REMOVED)
L46 70 S L45 AND MUTAT?
L47 27 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(560 OR 561 OR 562
L48 11 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(566 OR 567 OR 568
L49 37 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(841 OR 842 OR 843
L50 16 S L27 AND ((MUTAT? OR CODON OR RESIDUE)(30A)(845 OR 846 OR 847
L51 60 S L47 OR L48 OR L49 OR L50
L52 20 DUP REM L51 (40 DUPLICATES REMOVED)
L53 0 S L52 AND PY<=2002
L54 2 S L52 AND PY<=2003
L55 349 S L27 AND ((EXON(1A)10)OR(EXON(1A)12))

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L56      111 DUP REM L55 (238 DUPLICATES REMOVED)
L57      1 S L56 AND PY<=2002
L58      186 S L27 AND ((EXON(1A)18))
L59      67 DUP REM L58 (119 DUPLICATES REMOVED)
L60      0 S L59 AND PY<=2002
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=> log y